

Exhibit 1

Blast Result derived aligned sequence with that of Cat pKVC103rB sequence

>gi|4557788|ref|NM_000266.1| Homo sapiens Norrie disease (pseudoglioma)
(NDP), mRNA

Length = 1846

Score = 133 bits (67), Expect = 1e-28
Identities = 137/154 (88%), Gaps = 5/154 (3%)
Strand = Plus / Plus

Query: 2 gctgcatctttttctatgctctccctgctggcgctgatgggagatacagacagcaaaacg 61
||||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||
Sbjct: 427 gctgcatccttttctatgctctccctgctgggtgataatgggagatacagacagtaaaacg 486

Query: 62 gacagctcgtntcatgatcgactcggaccctctgcgncctgcatgaggcaccactatgttg 121
||||||| | |||| | ||||||| ||||| ||||||| ||||||| ||||||| ||
Sbjct: 487 gacagctcat-tcataatggactcggaccctcgacg-ctgcatgaggcaccactatg-tg 543

Query: 122 nattctatcagctcacccgttgntacaagtgtag 155
||||||| |||||| ||| |||||||
Sbjct: 544 gattctatcag-tcacccattg-tacaagtgtag 575

>gi|35016|emb|X65882.1|HSNDPG H.sapiens mRNA NDP
Length = 1846

Score = 133 bits (67), Expect = 1e-28
Identities = 137/154 (88%), Gaps = 5/154 (3%)
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Query: 2 gctgcatctttttctatgctctccctgctggcgctgatgggagatacagacagcaaaacg 61
||||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||
Sbjct: 427 gctgcatccttttctatgctctccctgctgggtgataatgggagatacagacagtaaaacg 486

Query: 62 gacagctcgtntcatgatcgactcggaccctctgcgncctgcatgaggcaccactatgttg 121
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Sbjct: 487 gacagctcat-tcataatggactcggaccctcgacg-ctgcatgaggcaccactatg-tg 543

Query: 122 nattctatcagctcacccgttgntacaagtgtag 155
||||||| |||||| ||| |||||||
Sbjct: 544 gattctatcag-tcacccattg-tacaagtgtag 575